

INFORMAL SEQUENCE LISTING

SEQ ID NO:1

Wild type DT with signal sequence underlined

5 GTGAGCAGAAACTGTTTGCGTCAATCTTAATAGGGGCGCTACTGGGGATAGGGGCCCCACCTTCAGCCCATGCA
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SEQ ID NO:2

30 DT-GML1

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SEQ ID NO: 3

DT-GML2

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SEQ ID NO: 4

DT-GMU2

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40 TGGAAGGGTTTTATAGTACCGACAATAAATACGACGCTGCGGGATACTCTGTAGATAATGAAAACCCGCTCTCT
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SEQ ID NO:5

DT-GMU3

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25 SEQ ID NO:6

DT-EGFL1

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30 GGAAAAGCTGGAGGCGTGGTCAAAGTGACGTATCCAGGACTGACGAAGGTTCTCGCACTAAAAGTGGATAATGCC
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5 SEQ ID NO:7

DT-EGFL2

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SEQ ID NO:8

DT-EGFU2

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SEQ ID NO:9

DT-EGFU3

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10 TGGAAAGGGTTTTATAGTACCGACAATAAATACGACGCTGCGGGATACTCTGTAGATAATGAAAACCCGCTCTCT
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20 ACAACTGCTGCTCTTTTCGATACTTCCTGGTATCGGTAGCGTAATGGGCATTGCAGACGGTGCCGTTACCCACAAT
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25 GTTGGCTACATCGGTGAGCGCTGTCAAGTATCGAGATCTGAAATGGTGGAACCTTAGATAA

SEQ ID NO:10

DT-IL2L1

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35 ATTAATAACTGGGAACAGGCGAAAGCGTTAAGCGTAGAACTTGAGATTAATTTTGAAACCCGTGGAAAACGTGGC
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5 AATCCCAAACCTACCAGGATGCTCACATTTAAGTTTTACATGCCCAAGAAGGCCACAGAAGTGAACATCTTCAG
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ACTTGA

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SEQ ID NO:11

DT-IL2L2

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15 TGGAAAGGGTTTTATAGTACCGACAATAAATACGACGCTGCGGGATACTCTGTAGATAATGAAAACCCGCTCTCT
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ACTTGA

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SEQ ID NO:12

DT-IL2U2

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40 TGGAAAGGGTTTTATAGTACCGACAATAAATACGACGCTGCGGGATACTCTGTAGATAATGAAAACCCGCTCTCT
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5 TCATGCATAAATCTTGATTGGGATGTCATAAGGGATAAACTAAGACAAAGATAGAGTCTTTGAAAGAGCATGGC
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15 GAAGAAGAACTCAAACCTCTGGAGGAAGTGCTAAATTTAGCTCAAAGCAAAAACCTTCACTTAAGACCCAGGGAC
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SEQ ID NO:13

20 DT-IL2U3
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25 GAACTATTAAGAAAGAGTTAGGTTTAAAGTCTCACTGAACCGTTGATGGAGCAAGTCGGAACGGAAGAGTTTATC
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40 TTAATCAGCAATATCAACGTAATAGTTCTGGAAGTAAAGGGATCTGAAACAACATTTCATGTGTGAATATGCTGAT
GAGACAGCAACCATTGTAGAATTTCTGAACAGATGGATTACCTTTTGTCAAAGCATCATCTCAAACTGACTTGA

SEQ ID NO:14

T7 promoter primer (5' primer for DT constructs)

5 GTAATACGACTCACTATAGGGC

SEQ ID NO:15

3' mutagenic primer for U2 constructs

10 GATTTATGCATGACAATGAGCTACCTGCTGATCTTCCACTTCCATTTCTGCACAGGCTTG

SEQ ID NO:16

3' mutagenic primer for U3 constructs

15 GATTTATGCATGACAATGAGCTACCTGCTGATTTTCCACTTCCATTTCTGCACAGGCTTG

SEQ ID NO:17

3' mutagenic primer for L1 constructs

20 GATTTATGCATGACAATGAGCTACCTTGACTCAACATTCCTAATGGTCCATTTCTGCACAGGCTTG

SEQ ID NO:18

3' mutagenic primer for L2 constructs

25 GATTTATGCATGACAATGAGCTACCTTGTGCCATAATCCTAATGGTCCATTTCTGCACAGGCTTG

SEQ ID NO:19

MMP substrate octapeptide for L1 constructs

30 GPLGMLSQ

SEQ ID NO:20

MMP substrate octapeptide for L2 constructs

35 GPLGLWAQ

SEQ ID NO:21

uPA substrate hexapeptide for U2 constructs

40 GSGRSA

SEQ ID NO:22

uPA substrate hexapeptide for U3 constructs

45 GSGKSA